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A-345C

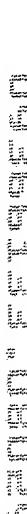
## **DIVISION - CONTINUATION - CONTINUATION-IN-PART APPLICATION TRANSMITTAL FORM**

Prior Application: Anticinated Classification Of This Application:

Class 435	Subcl	ass E	Examiner G. Draper			60				
To the Assistant Commissioner for Patents:										
This is a request for filing a  continuation  civisional  continuation-in-part application, under 37 CFR 1.53(b), of pending prior application Serial No. 08/920,608  filed on  August 27  , 19 97 ,  of  Pelleymounter, et al.  OB PROTEIN COMPOSITIONS AND METHODS  For CONTINUATION or DIVISIONAL APPLNs only: The entire disclosure of the prior application, from which an oath or declaration is supplied under Box 1b, below, is considered a part of the disclosure of the accompanying continuation or divisional application and is hereby incorporated by reference. The incorporation can only be relied upon when a portion has been inadvertently omitted from the submitted application parts.  1.  Transmitted herewith are:										
<ul> <li>✓ 4 pages of Sequence Listing; sequence statement.</li> <li>2. ✓ The filing fee is calculated below:</li> </ul>										
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3. ☑ Please charge Deposit Account No. 01-0519, in the name of Amgen Inc., in the amount of \$838.00. An original and one copy are enclosed.  4. ☑ Throughout the prosecution of this application, if any extension of time is necessary, please consider this a request therefor.  5. ☑ The Commissioner is hereby authorized to charge any additional filing fees which may be required by the accompanying application, any additional fees which may be required during pendency of this application as required by 37 CFR 1.16 or 1.17, or credit any overpayment to Deposit Account No. 01-0519 throughout the prosecution of this application.  6. ☐ Cancel in this application original claims of the prior application before calculating the filing fee. (At least one original independent claim must be retained for filing purposes.)										
	continuation [Serial No. 08/920,608]  DSITIONS AND METH NAL APPLNs only: The elered a part of the disclosure of specification, 2 of drawings.  Of Oath or Declaration by executed (original or y from a prior application of Sequence Listing; seculated below:    Number Filed   12	ner for Patents:    continuation	mer for Patents:    Continuation	ner for Patents:    Continuation	ner for Patents:    Continuation	Class 435  Subclass Examiner G. Draper Art Unit \$\frac{1}{1646}\$.  There for Patents:  Continuation divisional continuation-in-part application, under 37 CFR 1.53(b), Serial No. 08/920,608 filed on August 27 , 19 97 ,  DITIONS AND METHODS  NAL APPLNs only: The entire disclosure of the prior application, from which an oath or declaration is supplied ared a part of the disclosure of the accompanying continuation or divisional application and is hereby incorporated in can only be relied upon when a portion has been inadvertently omitted from the submitted application parts. the are:  of specification, 2 pages of claim(s) and 1 page of abstract, totaling 30 pages.  of of drawings.  of Oath or Declaration by the applicanti(s):  ty executed (original or copy)  y from a prior application (37 CFR 1.63(d)) (for continuation/divisional applies. only)  of Sequence Listing; sequence statement.  culated below:  Number Filed Number Extra Rate Fee 12 - 20 = 0 x \$18.00 = \$0.00				

# **EXPRESS MAIL CERTIFICATE**

"Express Mail" mail labeling number	EL198797275US	Date of Deposit.	August 2, 1999		
I hereby certify that this paper or fee is t	- ·	•		service under 37 CFR 1	10 on the dat
indicated above and is addressed to Bo	x Patent Application, Assistant Com	nissioner for Patents, Washington, D	OC 20231		
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# IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

Applicants: PELLEYMOUNTER, et al.

Serial No.: Not Yet Received Group Art Unit No.: Not Yet Assigned

Filed: August 2, 1999 Examiner: Not Yet Assigned

For: OB PROTEIN COMPOSITIONS AND METHODS

Docket No.: A-345C

### **PRELIMINARY AMENDMENT UNDER 37 CFR 1.121**

Assistant Commissioner for Patents Washington, DC 20231

Sir:

Please amend the accompanying application under 37 CFR 1.12 as follows:

#### In the Specification

Please amend the specification by inserting as the first paragraph the following:

--This application is a continuation of application Serial No. 08/920,608, filed August 27, 1997; which is a continuation of 08/474,833, filed June 7, 1995, now abandoned, which are hereby incorporated by reference.--

At page 3, line 29: after "SEQ. ID. Nos." delete --2 and 4-- and insert --3 and 6--

At page 11, line 17: delete "SEQ. ID. No. 2"

At page 11, line 19: delete "SEQ. ID. No. 1"

At page 12, line 4: after "3," delete --and 4-- and insert --4, 5, and 6--

At page 12, line 8: after "SEQ. ID. No." delete --3-- and insert --4--

At page 13, line 7: delete "SEQ. ID. No. 1 or 2"

At page 13, line 9: delete "SEQ. ID. No. 1"

At page 16, line 2: after "SEQ. ID. Nos." delete --1 and 2-- and insert --1, 2 and 3--

At page 25, line 2: after "SEQ. ID. Nos." delete --3 and 4-- and insert --4, 5 and 6--

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Freddie Craft

Printed Name

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### In the Claims

- 1. (Once amended) A method of treating excess weight in a mammal by continuous administration of 1 mg protein/kg body weight/day or less of an OB protein selected from the group consisting of:
  - (a) recombinant methionyl murine OB protein [SEQ. ID. No. 2)];
  - (b) recombinant methionyl human OB protein [(SEQ ID No. 1)];
  - (c) the protein of (a) or (b) lacking the methionyl residue at position -1;
  - (d) the protein of (a), (b) or (c) lacking a glutamine at position 28; and
  - (e) a chemically modified derivative of (a), (b),(c) or (d).
  - 8. (Once amended) A DNA sequence according to SEQ ID No. [3] 4.
- 11. (Once amended) A method of refolding partially purified OB protein in a solution obtained from inclusion bodies, said partially purified OB protein selected from the group consisting of:
  - (a) recombinant methionyl murine OB protein (SEQ. ID. No. [2] 3);
  - (b) recombinant methionyl human OB protein (SEQ ID No. [1] 6);
  - (c) the protein of (a) or (b) lacking the methionyl residue at position -1; wherein said refolding is accomplished using N-lauroyl sarcosine.

Please charge any fees associated herewith to Deposit Account 01-0519 and consider this a petition therefore if appropriate.

Respectfully submitted,

Karol M. Pessin

Attorney for Applicants

Registration No.: 34,899 Phone: (805) 447-2193 Date: August 2, 1999

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#### OB PROTEIN COMPOSITIONS AND METHOD

#### Field of the Invention

The present invention relates to OB protein compositions and methods for preparation and use thereof.

#### Background

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Although the molecular basis for obesity is 10 largely unknown, the identification of the "OB gene" and protein encoded by ("OB protein") has shed some light on mechanisms the body uses to regulate body fat deposition. Zhang et al., Nature 372: 425-432 (1994); see also, the Correction at Nature 374: 479 (1995). 15 The OB protein has been demonstrated to be active in vivo in both ob/ob mutant mice (mice obese due to a defect in the production of the OB gene product) as well as in normal, wild type mice. The biological activity manifests itself in, among other things, 20 weight loss. To date, however, optimum conditions for obtaining the rapid weight loss in normal animals has not been ascertained. In fact, some studies have shown that, when administered by injection, rather large dosages (10 mg of recombinant murine protein/kg body 25 weight/day) are necessary for normal mice to lose 2.6% of their body weight (at the end of a 32 day period). While presently uncertain, one explanation for the necessity of such large dosages is that the optimum weight loss effects are seen predominantly when the 30 protein is in constant circulation, a condition that may not be efficiently achieved by injecting the protein.

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#### Summary of the Invention

The present invention stems from the observation that, as compared to administering OB protein by injection, administering OB protein by continuous pump infusion results in equivalent (or better) weight loss, in a shorter time, and with substantially lower dosages. The working example below demonstrates that a dose of 0.5 mg protein/kg body weight/day, administered via implantable osmotic pump, results in a weight loss of over 4% (as compared to baseline weight). This is in substantial contrast to other studies where similar, or less weight loss (at a comparable time point) was observed with intraperitoneal injection at the relatively high dosage of 10 mg of protein/kg body weight/day.

Thus, one aspect of the present invention is a method of treating excess weight by administering OB protein in a form for constant supply, at a dosage of less than or equal to about 1 mg protein/kg body weight/day. The dosage of less than or equal to about 1 mg protein/kg/day refers to dosages sufficient to result in observable weight loss. This is apparent from the present studies where a dosage of 0.5 mg/kg/day was sufficient to result in observable weight loss when continuously administered. In studies where injection had been the mode of administration, far higher dosages were required for weight loss. At injection dosages of 0.1 and 1 mg/kg/day, substantially no weight loss was observed in wild type (normal) mice. For example, in one study, at a comparable time point (6th day), there was a .2% loss at the 1 mg/kg dose (data not shown). Minimal weight loss was observed at the relatively high 10 mg/kg/day dose. (1.9% weight loss at day 6, data not shown). Thus, the present invention provides for dosages of 1 mg/kg/day or less

when administered so that the supply of protein is continuous.

Connected with the present studies are the compositions and methods used for production of recombinant murine and human OB protein. The first example below discloses the preparation of recombinant murine protein, and the second example below discloses the preparation of recombinant human protein.

Additional aspects of the present invention, therefore, include the below compositions and methods for preparing recombinant murine methionyl OB protein and recombinant human methionyl OB protein, including DNA sequences, vectors, host cells, methods of fermentation, and methods of purification.

Detailed Description

The present invention stems from the observation that continuous administration of OB protein results in the need for much lower dosages for weight loss than those dosages required by acute daily injection. As set forth above, a dosage of 1 mg protein/kg body weight/day or less, continuously administered, resulted in rapid weight loss. When the underivatized protein was administered by acute injection at the 1 mg/kg/day dose, almost no weight loss in wild type (normal) mice.

The OB protein may be selected from the recombinant murine and human methionyl proteins set forth below (SEQ. ID Nos. 2 and 4) or those lacking a glutaminyl residue at position 28. (See Zhang et al, Nature, supra, at page 428.) The recombinant human OB gene product is, as a mature protein, 146 amino acids; some of the DNAs obtained were observed to encode a protein lacking a glutamine residue at position 28. Zhang et al., Nature 372 at 428. The murine protein is substantially homologous to the human protein,

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particularly as a mature protein, and, further, particularly at the N-terminus. One may prepare an analog of the recombinant human protein by altering (such as substituting amino acid residues), in the recombinant human sequence, the amino acids which diverge from the murine sequence. Because the recombinant human protein has biological activity in mice, such analog would likely be active. Proteins lacking an N-terminal methionyl residue, such as those produced by eukaryotic expression, are also available for use.

In addition, although the present working example involved continuous administration via implantable pump, it is contemplated that other modes of continuous administration may be practiced. For example, chemical derivatization may result in sustained release forms of the protein which have the effect of continuous presence in the blood stream, in predictable amounts. Thus, one may derivatize the above proteins to effectuate such continuous administration. The dosage of 1 mg protein/kg body weight/day or less herein refers to the mass of protein, exclusive of other chemical moieties used to derivatize the protein.

Generally, the present protein (herein the term "protein" is used to include "peptide", unless otherwise indicated) may be derivatized by the attachment of one or more chemical moieties to the protein moiety. The chemically modified derivatives may be further formulated for intraarterial, intraperitoneal, intramuscular subcutaneous, intravenous, oral, nasal, pulmonary, topical or other routes of administration. Chemical modification of biologically active proteins has been found to provide additional advantages under certain circumstances, such as increasing the stability and circulation time of the

therapeutic protein and decreasing immunogenicity. U.S. Patent No. 4,179,337, Davis et al., issued December 18, 1979. For a review, see Abuchowski et al., in Enzymes as Drugs. (J.S. Holcerberg and J. Roberts, eds. pp. 367-383 (1981)). A review article describing protein modification and fusion proteins is Francis, Focus on Growth Factors 3: 4-10 (May 1992) (published by Mediscript, Mountview Court, Friern Barnet Lane, London N20, OLD, UK). For the present continuous administration, it is preferred that the 10 chemical modification allow for an increase in circulation time of the protein, so that a dosage of about 1 mg protein (exclusive of chemical moiety)/kg body weight of a mammal/day or less will result in weight loss of a mammal. The present continuous 15 administration will provide for weight loss of approximately 5% of body mass in a period of 7 or fewer days.

The chemical moieties suitable for derivatization may be selected from among water soluble 20 polymers. The polymer selected should be water soluble so that the protein to which it is attached does not precipitate in an aqueous environment, such as a physiological environment. Preferably, for therapeutic use of the end-product preparation, the polymer will be 25 pharmaceutically acceptable. One skilled in the art will be able to select the desired polymer based on such considerations as whether the polymer/protein conjugate will be used therapeutically, and if so, the desired dosage, circulation time, resistance to 30 proteolysis, and other considerations. For the present proteins and peptides, the effectiveness of the derivatization may be ascertained by administering the derivative, in the desired form (i.e., by osmotic pump, or, more preferably, by injection or infusion, or, 35

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further formulated for oral, pulmonary or nasal delivery, for example), and measuring weight loss.

The water soluble polymer may be selected from the group consisting of, for example, polyethylene glycol, copolymers of ethylene glycol/propylene glycol, carboxymethylcellulose, dextran, polyvinyl alcohol, polyvinyl pyrrolidone, poly-1, 3-dioxolane, poly-1,3,6-trioxane, ethylene/maleic anhydride copolymer, polyaminoacids (either homopolymers or random copolymers), and dextran or poly(n-vinyl pyrrolidone)polyethylene glycol, propropylene glycol homopolymers, prolypropylene oxide/ethylene oxide co-polymers, polyoxyethylated polyols and polyvinyl alcohol. Polyethylene glycol propionaldenhyde may have advantages in manufacturing due to its stability in water.

The polymer may be of any molecular weight, and may be branched or unbranched. For polyethylene glycol, the preferred molecular weight is between about 2kDa and about 100kDa (the term "about" indicating that in preparations of polyethylene glycol, some molecules will weigh more, some less, than the stated molecular weight) for ease in handling and manufacturing. Other sizes may be used, depending on the desired therapeutic profile (e.g., the duration of sustained release desired, the effects, if any on biological activity, the ease in handling, the degree or lack of antigenicity and other known effects of the polyethylene glycol to a therapeutic protein or analog).

The number of polymer molecules so attached may vary, and one skilled in the art will be able to ascertain the effect on function. One may monoderivatize, or may provide for a di-, tri-, tetra- or some combination of derivatization, with the same or different chemical moieties (e.g., polymers, such as

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different weights of polyethylene glycols). The proportion of polymer molecules to protein (or peptide) molecules will vary, as will their concentrations in the reaction mixture. In general, the optimum ratio (in terms of efficiency of reaction in that there is no excess unreacted protein or polymer) will be determined by factors such as the desired degree of derivatization (e.g., mono, di-, tri-, etc.), the molecular weight of the polymer selected, whether the polymer is branched or unbranched, and the reaction conditions.

The polyethylene glycol molecules (or other chemical moieties) should be attached to the protein with consideration of effects on functional or antigenic domains of the protein. There are a number of attachment methods available to those skilled in the art. E.g., EP 0 401 384 herein incorporated by reference (coupling PEG to G-CSF), see also Malik et al., Exp. Hematol. <u>20</u>: 1028-1035 (1992) (reporting pegylation of GM-CSF using tresyl chloride). For example, polyethylene glycol may be covalently bound through amino acid residues via a reactive group, such as, a free amino or carboxyl group. Reactive groups are those to which an activated polyethylene glycol molecule may be bound. The amino acid residues having a free amino group may include lysine residues and the N-terminal amino acid residue. Those having a free carboxyl group may include aspartic acid residues, glutamic acid residues, and the C-terminal amino acid Sulfhydrl groups may also be used as a residue. reactive group for attaching the polyethylene glycol molecule(s). Preferred for therapeutic purposes is attachment at an amino group, such as attachment at the N-terminus or lysine group. Attachment at residues important for receptor binding should be avoided if receptor binding is desired.

One may specifically desire N-terminally chemically modified protein. Using polyethylene glycol as an illustration of the present compositions, one may select from a variety of polyethylene glycol molecules (by molecular weight, branching, etc.), the proportion of polyethylene glycol molecules to protein (or peptide) molecules in the reaction mix, the type of pegylation reaction to be performed, and the method of obtaining the selected N-terminally pegylated protein.

The method of obtaining the N-terminally pegylated preparation (i.e., separating this moiety from other monopegylated moieties if necessary) may be by purification of the N-terminally pegylated material from a population of pegylated protein molecules.

Selective N-terminal chemical modification may be accomplished by reductive alkylation which exploits differential reactivity of different types of primary amino groups (lysine versus the N-terminal) available for derivatization in a particular protein. Under the

appropriate reaction conditions, substantially selective derivatization of the protein at the N-terminus with a carbonyl group containing polymer is achieved. For example, one may selectively N-terminally pegylate the protein by performing the

reaction at a pH which allows one to take advantage of the pKa differences between the  $\epsilon$ -amino group of the lysine residues and that of the  $\alpha$ -amino group of the N-terminal residue of the protein. By such selective derivatization, attachment of a water soluble polymer

to a protein is controlled: the conjugation with the polymer takes place predominantly at the N-terminus of the protein and no significant modification of other reactive groups, such as the lysine side chain amino groups, occurs. Using reductive alkylation, the water

35 soluble polymer may be of the type described above, and should have a single reactive aldehyde for coupling to

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the protein. Polyethylene glycol propionaldehyde, containing a single reactive aldehyde, may be used.

In yet another aspect of the present invention, provided are methods of using pharmaceutical compositions of the proteins and derivatives. Such pharmaceutical compositions may be for administration for injection, or for oral, pulmonary, nasal or other forms of administration which allow for the desired circulating dose of about 1 mg protein/kg body weight/day or less. In general, comprehended by the invention are pharmaceutical compositions comprising effective amounts of protein or derivative products of the invention together with pharmaceutically acceptable diluents, preservatives, solubilizers, emulsifiers, adjuvants and/or carriers. Such compositions include diluents of various buffer content (e.g., Tris-HCl, acetate, phosphate), pH and ionic strength; additives such as detergents and solubilizing agents (e.g., Tween 80, Polysorbate 80), anti-oxidants (e.g., ascorbic acid, sodium metabisulfite), preservatives (e.g., Thimersol, benzyl alcohol) and bulking substances (e.g., lactose, mannitol); incorporation of the material into particulate preparations of polymeric compounds such as polylactic acid, polyglycolic acid, etc. or into liposomes. Hylauronic acid may also be used, and this may have the effect of promoting sustained duration in the circulation. Such compositions may influence the physical state, stability, rate of in vivo release, and rate of in vivo clearance of the present proteins and derivatives. See, e.g., Remington's Pharmaceutical Sciences, 18th Ed. (1990, Mack Publishing Co., Easton, PA 18042) pages 1435-1712 which are herein incorporated by reference. The compositions may be prepared in liquid form, or may be in dried powder, such as lyophilized form. The

effective amounts are those herein described.

The OB proteins and derivatives described are useful for modulation of the rate or quantity of fat cell deposition in a mammal. This is thought to be accomplished, in part, by a reduction in appetite,

- i.e., a reduction in food intake. Thus, one observable result is weight loss, or, put another way, a method of treating excess weight (via weight loss). Thus, the present compositions are useful for the manufacture of a medicament for treating excess weight in a mammal.
- Another aspect is a method for reducing appetite.

  Either of these aspects, modulation of fat deposition or modulation of appetite, are particularly important treatments for humans (or other mammals) who desire to lose weight.
- One skilled in the art will be able to ascertain other effective dosages by administration and observing weight loss. Here, the dosage of 1 mg protein/kg body weight/day or less was seen to be particularly effective, when administered on a
- continuous basis. More particularly, the dosage of 0.5 mg/kg body weight/day was seen to be particularly effective on normal mice. Excess weight refers to body mass for which removal is desired. It is contemplated that the present compositions and methods will be used
- to treat cases where removal of such excess weight (as a result of the present invention) will benefit other health concerns, such as diabetes, high blood pressure or cardiac problems, high cholesterol levels, low locomotion levels and other manifestations of excess
- weight. As such, the present compositions and methods may be used in conjunction with other medicaments, such as those useful for the treatment of diabetes (e.g., insulin, and possibly amylin), cholesterol and blood pressure lowering medicaments, and locomotion
- increasing medicaments (e.g., amphetamines). Such

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administration may be simultaneous or may be <u>in</u> <u>serriatim</u>.

In addition, the present compositions and methods may be used in conjunction with surgical procedures, such as cosmetic surgeries designed to alter the overall appearance of a body (e.g., liposuction or laser surgeries designed to reduce body mass). The health benefits of cardiac surgeries may be increased with concomitant use of the present compositions and methods.

Therefore, the present invention encompasses a method of treating excess weight in a mammal by continuous administration of 1 mg protein/kg body weight/day or less of an OB protein selected from the group consisting of:

- (a) recombinant methionyl murine OB protein (SEQ. ID. No. 2);
- (b) recombinant methionyl human OB protein (SEQ ID No. 1);
- (c) the protein of (a) or (b) lacking the methionyl residue at position -1;
  - (d) the protein of (a), (b) or (c) lacking a glutamine at position 28; and
- (e) a chemically modified derivative of (a),(b),(c) or (d), wherein the chemical modificationallows for an increase in circulation time.

Preferably, the composition of subpart (e) is a pegylated derivative, and, more preferably, an N-terminally pegylated derivative.

- The derivative of subpart (e) allows for continuous administration of the protein by increasing the circulation time of the (unmodified) protein. The present invention also encompasses a method of treating excess weight where the method of continuous
- 35 administration is by implantable pump, such as an osmotic pump.

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In other aspects, the present invention relates to recombinant murine and recombinant human OB DNAs and proteins, such as those of SEQ. ID NOs. 1, 2, 3, and 4, below. The recombinant proteins below are bacterially expressed, and contain N-terminal methionyl residues. Vectors and host cells useful for producing such proteins are also provided. The vectors include pCFM1656 containing SEQ ID No. 1 or 3, and host cells containing such vectors.

Methods for preparation of the recombinant proteins are also provided, including methods for fermentation and methods for purification.

In particular, the use of sarcosine for refolding of OB protein in solution, obtained from bacterial inclusion bodies, provided for extremely efficient refolding. When proteins are expressed in bacteria, they may not be in the proper threedimensional configuration, or, as referred to herein, properly refolded. The three dimensional configuration may be critical for biological activity, and storage stability. Although Sarckosyl has been used in processes for purification of another protein (G-CSF, e.g., WO 89/10932), surprisingly, the use of sarcosine for the OB protein has resulted in a refolding efficiency of over 95%. Contemplated herein is the use of N-lauroylsarcosine in a range of 0.5% - 2.0 % weight per volume of OB protein in solution (obtained from inclusion bodies). With the use of 1% sodium sarcosine, the refolding efficiency, as determined by SDS PAGE and reverse phase HPLC, was 95% or greater. While one skilled in the art will recognize that other compositions may be used for refolding, the use of N-lauroyl sarcosine, as illustrated in the examples below, is particularly advantageous for providing extremely efficient refolding. The removal of

sarcosine was accomplished using Dowex®.

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Therefore, the present invention also includes a method of refolding partially purified OB protein in a solution obtained from inclusion bodies, said partially purified OB protein selected from the group consisting of:

- (a) recombinant methionyl murine OB protein (SEQ. ID. No. 2);
- (b) recombinant methionyl human OB protein (SEO ID No. 1);
- (c) the protein of (a) or (b) lacking the methionyl residue at position -1;

wherein said refolding is accomplished using sarcosine.

The present invention also includes methods

of wherein said N-lauroyl sarcosine is used at a

concentration of 0.5% - 2.0% weight per volume of

solution, and, more particularly, the use of 1% N
lauroyl sarcosine. An oxidizing agent, such as copper

sulfate, is also used in the refolding process.

The following examples are offered to more fully illustrate the invention, but are not to be construed as limiting the scope thereof.

# EXAMPLE 1: <u>Use of Murine OB Protein in a</u> <u>Continuous Pump Infusion System</u>

This example demonstrates that continuous infusion of OB protein results in weight loss in normal mice. Normal (non-obese) mice were administered murine OB protein via osmotic pump infusion. A dosage of 0.5 mg protein/kg body weight/day resulted in a 4.62% (+/-1.34%) loss from baseline weight by the 6th day of infusion.

#### MATERIALS AND METHODS

Animals: Wild type (+/+) C57B16 mice were used for this experiment. The age of the mice at the initial time point was 8 weeks, and the animals were weight stabilized. 10 mice were used for each cohort (vehicle vs. protein).

#### Animal Handling.

10 Feeding and weight measurement. Mice were given ground rodent chow (PMI Feeds, Inc.) in powdered food feeders (Allentown Caging and Equipment) which allowed a more accurate and sensitive measurement than use of regular block chow. Weight was measured at the same time each day (2:00 p.m.), for a period of 6 days. Body weight on the day prior to the infusion was defined as baseline weight. The mice used weighed 18-22 grams.

Housing. Mice were single-housed, and maintained under humane conditions.

Administration of Protein or Vehicle. Protein (as described below) or vehicle (phosphate buffered saline, ph 7.4) were administered by osmotic pump infusion. Alzet osmotic minipumps (Alza, Palo Alto, CA, model no. 1007D) were surgically placed in each mice in a subcutaneous pocket in the subscapular area. The pumps were calibrated to administer 0.5 µl protein in solution per hour for a dosage of 0.5 mg protein/kg body weight/day.

<u>Controls</u>: Control animals were those who had a Alzet osmotic minipump infusing phosphate buffered saline (pH 7.4).

Protein: Recombinant murine OB protein was used for the present experiments, generally at a concentration of about 0.9 mg/ml phosphate buffered saline, pH 7.4. The amino acid sequence (and DNA sequence) used was the following:

# Recombinant murine met OB (double stranded) DNA and amino acid sequence (Seq. ID. Nos. 1 and 2):

C.	^	TCTAGATTTGAGTTTTAACTTTTAGAAGGAGGAATAACATATGGTACCGATCCAGAAAGT									
5	9	AGATCTAAACTCAAAATTGAAAATCTTCCTCCTTATTGTATACCATGGCTAGGTCTTTCA  M V P I Q K V	-								
10 69	69	TCAGGACGACACCAAAACCTTAATTAAAACGATCGTTACGCGTATCAACGACATCAGTCA	128								
_ •		AGTCCTGCTGTGGTTTTGGAATTAATTTTGCTAGCAATGCGCATAGTTGCTGTAGTCAGT Q D D T K T L I K T I V T R I N D I S H	-								
15	129	CACCCAGTCGGTCTCCGCTAAACAGCGTGTTACCGGTCTGGACTTCATCCCGGGTCTGCA	188								
		GTGGGTCAGCCAGAGGCGATTTGTCGCACAATGGCCAGACCTGAAGTAGGGCCCAGACGT T Q S V S A K Q R V T G L D F I P G L H	-								
20	189	<del></del>	248								
		GGGCTAGGATTCGAACAGGTTTTACCTGGTCTGGGACCGACATATGGTCGTCCACAATTG PILS K M D Q T L A V Y Q Q V L T	-								
25	249	- <del></del>	308								
		GAGGGACGGCAGGGTCTTGCAAGAAGTCTAGCGATTGCTGGAGCTCTTGGAAGCGCTGGA S L P S Q N V L Q I A N D L E N L R D L	-								
30	309	GCTGCACCTGCTGGCATTCTCCAAATCCTGCTCCCTGCCGCAGACCTCAGGTCTTCAGAA	368								
		CGACGTGGACGACCGTAAGAGGTTTAGGACGAGGGACGGCGTCTGGAGTCCAGAAGTCTT  L H L L A F S K S C S L P Q T S G L Q K	-								
35 369	369	ACCGGAATCCCTGGACGGGTCCTGGAAGCATCCCTGTACAGCACCGAAGTTGTTGCTCT	428								
		TGGCCTTAGGGACCTGCCCCAGGACCTTCGTAGGGACATGTCGTGGCTTCAACAACGAGA PESLDGVLEASLYSTEVVALL	-								
40	429	GTCCCGTCTGCAGGGTTCCCTTCAGGACATCCTTCAGCAGCTGGACGTTTCTCCGGAATG	488								
		CAGGGCAGACGTCCCAAGGGAAGTCCTGTAGGAAGTCGTCGACCTGCAAAGAGGCCTTAC S R L Q G S L Q D I L Q Q L D V S P E C	-								
45	489	TTAATGGATCC -+ AATTACCTAGG									

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Herein, the first amino acid of the amino acid sequence for recombinant protein is referred to as +1, and is valine, and the amino acid at position -1 is methionine. The C-terminal amino acid is number 146 (cysteine).

The cloning of the murine OB DNA for expression in  $\underline{E}$ .  $\underline{coli}$  was done as follows. The DNA sequence was deduced from the published peptide sequence that appeared in Zhang et al., Nature 372:425-432 It was reverse translated using E. coli optimal (1994).10 The terminal cloning sites were XbaI to BamHI. codons. A ribosomal binding enhancer and a strong ribosomal binding site were included in front of the coding region. The duplex DNA sequence was synthesized using standard techniques. Correct clones were confirmed by 15 demonstrating expression of the recombinant protein and presence of the correct OB DNA sequence in the resident plasmid.

#### Expression Vector and Host Strain 20

The plasmid expression vector used was pCFM1656, ATCC Accession No. 69576. The above DNA was ligated into the expression vector pCFM1656 which had been linearized with XbaI and BamHI and transformed into the <u>E. coli</u> host strain, FM5. <u>E. coli</u> FM5 cells were derived at Amgen Inc., Thousand Oaks, CA from E. coli K-12 strain (Bachmann, et al., Bacteriol. Rev. 40: 116-167 (1976)) and contain the integrated lambda phage repressor gene, cI<sub>857</sub> (Sussman et al., C.R. Acad. Sci.

254: 1517-1579 (1962)). Vector production, cell 30 transformation, and colony selection were performed by standard methods. <u>E.g.</u>, Sambrook, et al., Molecular Cloning: A Laboratory Manual, 2d Edition, Cold Spring Harbor Laboratory Press, Cold Spring Harbor, NY. Host cells were grown in LB media.

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Fermentation Process A three-phase fermentation protocol was used known as a fed-batch process. Media compositions are set forth below.

Batch: A nitrogen and phosphate source were sterilized (by raising to 122 °C for 35 minutes, 18-20 psi) in the fermentation vessel (Biolafitte, 12 liter capacity). Upon cooling, carbon, magnesium, vitamin, and trace metal sources were added aseptically. An overnight culture of the above recombinant murine protein-producing bacteria (16 hours or more) of 500 mL (grown in LB broth) was added to the fermentor.

Feed I: Upon reaching between 4.0-6.0 OD<sub>600</sub>, cultures were fed with Feed I. The glucose was fed at a limiting rate in order to control the growth rate ( $\mu$ ). An automated system (called the Distributive Control System) was instructed to control the growth rate to 0.15 generations per hour.

Feed II: When the OD600 had reached 30, culture temperature was slowly increased to 42°C and the feed was changed to Feed II, below. The fermentation was then allowed to continue for 10 hours with sampling every 2 hours. After 10 hours, the contents of the fermentor was chilled to below 20°C and harvested by centrifugation.

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## Media Composition:

	Batch:	10 g/L	Yeast extract
		5.25  g/L	$(NH_4)_2SO_4$
		3.5 g/L	K <sub>2</sub> HPO <sub>4</sub>
5		4.0 g/L	KH <sub>2</sub> PO <sub>4</sub>
		5.0 g/L	Glucose
		1.0 g/L	$MgSO_4 \cdot 7H_2O$
		2.0 mL/L	Vitamin Solution
		2.0~mL/L	Trace Metal Solution
10		1.0 mL/L	P2000 Antifoam
	Feed I:	50 g/L	Bacto-tryptone
		50 g/L	Yeast extract
		450  g/L	Glucose
		8.75  g/L	$MgSO_4 \cdot 7H_2O$
15		10 mL/L	Vitamin Solution
		10  mL/L	Trace Metal Solution
	Feed II:	200 g/L	Bacto-tryptone
		100 g/L	Yeast extract
		110 g/L	Glucose
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Vitamin Solution (Batch and Feed I):

0.5 g Biotin, 0.4 g Folic acid, and 4.2 g riboflavin, were dissolved in 450 mls H<sub>2</sub>O and 3 mls 10 N NaOH, and brought to 500 mls in H<sub>2</sub>O. 14 g pyridoxine-HCl and 61 g niacin were dissolved 150 ml H<sub>2</sub>O and 50 ml 10 N NaOH, and brought to 250 ml in H<sub>2</sub>O. 54 g pantothenic acid was dissolved in 200 ml H<sub>2</sub>O, and brought to 250 ml. The three solutions were combined and brought to 10 liters total volume.

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Trace Metal Solution (Batch and Feed I):
Ferric Chloride (FeCl<sub>3</sub>·6H<sub>2</sub>O): 27 g/L

Zinc Chloride (ZnCl<sub>2</sub>·4H<sub>2</sub>O): 2 g/L

Cobalt Chloride (CoCl<sub>2</sub>·6H<sub>2</sub>O): 2 g/L

Sodium Molybdate (NaMoO<sub>4</sub>·2H<sub>2</sub>O): 2 g/L

Calcium Chloride (CaCl<sub>2</sub>·2H<sub>2</sub>O): 1 g/L

Cupric Sulfate (CuSO<sub>4</sub>·5H<sub>2</sub>O): 1.9 g/L

Boric Acid (H<sub>3</sub>BO<sub>3</sub>): 0.5 g/L

Manganese Chloride (MnCl<sub>2</sub>·4H<sub>2</sub>O): 1.6 g/L

Sodium Citrate dihydrate: 73.5 g/L

Purification Process for Murine OB Protein

Purification was accomplished by the

following steps (unless otherwise noted, the following steps were performed at 4°C):

- 1. Cell paste. <u>E</u>. <u>coli</u> cell paste was suspended in 5 times volume of 7 mM of EDTA, pH 7.0. The cells in the EDTA were further broken by two passes through a microfluidizer. The broken cells were centrifuged at 4.2 K rpm for 1 hour in a Beckman J6-B centrifuge with a JS-4.2 rotor.
- 2. Inclusion body wash #1. The supernatant from above was removed, and the pellet was resuspended with 5 times volume of 7 mM EDTA, pH 7.0, and homogenized. This mixture was centrifuged as in step 1.
  - 3. Inclusion body wash #2. The supernatant from above was removed, and the pellet was resuspended in ten times volume of 20 mM tris, pH 8.5, 10 mM DTT, and 1% deoxycholate, and homogenized. This mixture was centrifuged as in step 1.
  - 4. Inclusion body wash #3. The supernatant from above was removed and the pellet was resuspended in ten times volume of distilled water, and homogenized. This mixture was centrifuged as in step 1.

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- 5. Refolding. The pellet was refolded with 15 volumes of 10 mM HEPES, pH 8.5, 1% sodium sarcosine (N-lauroyl sarcosine), at room temperature. After 60 minutes, the solution is made to be 60  $\mu$ M copper sulfate, and then stirred overnight.
- 6. Removal of sarcosine. The refolding mixture was diluted with 5 volumes of 10 mM tris buffer, pH 7.5, and centrifuged as in step 1. The supernatant was collected, and mixed with agitation for one hour with Dowex® 1-X4 resin (Dow Chemical Co., Midland MI), 20-50 mesh, chloride form, at 0.066% total volume of diluted refolding mix. See WO 89/10932 at page 26 for more information on Dowex®. This mixture was poured into a column and the eluant was collected. Removal of sarcosine was ascertained by reverse phase HPLC.
  - 7. Acid precipitation. The eluant from the previous step was collected, and pH adjusted to pH 5.5, and incubated for 30 minutes at room temperature. This mixture was centrifuged as in step 1.
- 8. Cation exchange chromatography. The pH of the supernatant from the previous step was adjusted to pH 4.2, and loaded on CM Sepharose Fast Flow (at 7% volume). 20 column volumes of salt gradient were done at 20 mM NaOAC, pH 4.2, 0 M to 1.0 M NaCl.
- 9. Hydrophobic interaction chromatography. The CM Sepharose pool of peak fractions (ascertained from ultraviolet absorbance) from the above step was made to be 0.2 M ammonium sulfate. A 20 column volume reverse salt gradient was done at 5 mM NaOAC, pH 4.2, with .4 M to 0 M ammonium sulfate. This material was concentrated and diafiltered into PBS.

#### Results

Presented below are the percent (%) differences from baseline weight in C57B16J mice (8 weeks old):

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Table 1: Weight Loss Upon Continuous Infusion

Time (days)	<u>Vehicle (PBS)</u>	Recombinant OB <u>protein</u>
Days 1-2	3.24 +/- 1.13	1.68 +/- 1.4
Days 3-4	4.3 +/97	-2.12 +/79
Days 5-6	4.64 +/96	-4.62 +/- 1.3

As can be seen, at the end of a 6 day 10 continuous infusion regime, animals receiving the OB protein lost over 4% of their body weight, as compared to baseline. This is a substantially more rapid weight loss than has been observed with intraperitoneal (i.p.) injection. Weight loss at the end of a 32-day injection 15 period, in wild type (normal) mice, with daily i.p. injections of recombinant murine OB protein at a 10 mg/kg dose was 2.6%, and had not been more than 4% at any time during the dosing schedule (data not shown). The present data indicate that with continuous 20 infusion, a 20-fold lower dosage (0.5 mg/kg vs. 10 mg/kg) achieves more weight loss in a shorter time period.

The results seen here are statistically significant, e.g., -4.62% with p <.0001.

#### EXAMPLE 2: Dose Response Studies

An additional study demonstrated that there was a dose response to continuous administration of OB protein. In this study, non-obese, CD-1 mice, weighing 35-40 g were administered recombinant murine OB protein using methods similar to the above example. The results are set forth in Table 2, below, (with % body weight lost as compared to baseline, measured as above):

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Table 2: Dose Response With Continuous Administration

Dose	Time	% Reduction in
		body weight
0.03	Day 2	3.5
mg/kg/day		
1 mg/kg/day	Day 2	7.5
1 mg/kg/day	Day 4	14

As can be seen, increasing the dose from 0.03 mg/kg/day to 1 mg/kg/day increased the weight lost from 3.5% to 7.5%. It is also noteworthy that at day 4, the 1 mg/kg/day dosage resulted in a 14% reduction in body weight.

# 20 EXAMPLE 3: <u>Cloning and Expression of a</u> Recombinant Human Methionyl OB Protein

This example provides compositions and methods for preparation of a recombinant human version of the OB protein.

The human version of the OB DNA was constructed from the murine OB DNA, as in Example 1, above, by replacing the region between the MluI and BamHI sites with duplex DNA (made from synthetic oligonucleotides) in which 20 codon substitutions had been designed. The MluI site is shown

under the solid line in the sequence below. This DNA was put into the pCFM1656 vector (ATCC Accession No. 69576), in the same fashion as the recombinant murine protein, as described above. Herein, the first amino acid of the amino acid sequence for recombinant human protein below is referred to as +1, and is valine, and the amino acid at position -1 is methionine. The C-terminal amino acid is number 146 (cysteine).

Recombinant human met OB (Double Stranded) DNA and amino acid sequence (Seq. ID. Nos. 3 and 4)

_	4	CAT	TAT(	GGT	ACC(	GAT(	CCA	GAA	AGT'	TCA	GGA	CGA	CAC	CAA	AAC	CTT	\AT'	ΓΑΑ <i>Ι</i>	AAC(	GAT(	CGTT	60
5	1	GTA	ATA( M	CCA'	-+- TGG( P	CTA(	GGT( Q	+ CTT' K	rca V	AGT Q	CCT D					GAA'		ATT' K	rtg( T	CTA(	GCAA V	-
10	<i>C</i> 1				_											TAA 					AGGC	120
	61																				rccg	120
15		Т	R	Ι	N	D	I	S	Н	T	Q	S	V	S	S	K	Q	R	V	T	G	-
		CT	GGA	CTT	CAT	CCC	GGG'	TCT	GCA	CCC	GAT	CCT	GAC	CTT	GTC	CAA	AAT	GGA:	CCA	GAC(	CCTG	100
•	121																				+ GGAC	100
20		L	D	F	I	P	G	L	Н	P	I	L	${f T}$	Ŀ	S	K	M	D	Q	Т	L	-
25	101																				TAAC +	240
<i>4</i> . J	181																				ATTG	<b>4</b> ∓ <b>V</b>
		A	V	Y	Q	Q	I	L	Т	S	Μ	P	S	R	N	V	L	Q	I	S	N	-
30		GA(	CCT	CGA	GAA	CCT															CCTG	
	241		 GGA	 GCT	- + - CTT	 GGA															+ GGAC	300
35		D	L	E	N	L	R	D	L	L	Н	V	L	A	F	S	K	S	С	H	L	-
	201	CCATGGGCTTCAGGTCTTGAGACTCTGGACTCTCTGGGCGGGGTCCTGGAAGCATCCGGT											360									
40	301																				GCCA	300
		P	W	A	S	G	L	Ε	Т	L	D	S	Ŀ	G	G	V	L	E	A	S	G	-
45	361																				TTGG +	420
	301																				AACC	
50		Y	S	Т	Е	V	V	A	L	S	R	L	Q	G	S	Ŀ	Q	D	М	L	W	-
			CAGCTGGACCTGTCTCCGGGTTGTTAATGGATCC																			
	421		CGA	.CCI	- + - 'GGA	CAG	GAGG	+ CCC:						454	=							
55		Q	L	D	L	S	P	G	С	*												

<u>Fermentation</u>: Fermentation of the above host cells to produce recombinant human OB protein was accomplished using the conditions and compositions as described above

for recombinant murine material. The results were analyzed for yield (grams ob DNA product/liter of fermentation broth), prior to purification of the recombinant human OB material. (Minor amounts of bacterial protein were present.) Bacterial expression was also calculated.

Table 3: Analysis of Human OB Protein Expression

Timepoint	OD (@600 nm)	Yield (g/L)	Expression
			(mg/OD·L)
Ind. + 2 hours.	47	1.91	41
Ind. + 4 hours.	79	9.48	120
Ind. + 6 hours.	95	13.01	137
Ind. + 8 hours.	94	13.24	141
Ind. + 10 hours.	98	14.65	149

abbreviations: Ind. + \_\_\_ hours means the hours after induction of protein expression, as described in Example I for the recombinant murine material using pCFM1656

OD: optical density, as measured by spectrophotometer milligrams per OD unit per liter mg/OD·L: expression in terms of milligrams of protein per OD unit per liter.

g/L: grams protein/liter fermentation broth

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Purification of the recombinant human OB protein: Recombinant human protein may be purified using methods similar to those used for purification of recombinant murine protein, as in Example 1, above. For preparation of recombinant human OB protein, step 8 was performed by adjusting the pH of the supernatant from step 7 to pH 5.0, and loading this onto a CM Sepharose fast flow column. The 20 column volume salt gradient was performed at 20 mM NaOAC, pH 5.5, 0M to 0.5 M NaCl. Step 9 was performed by diluting the CM Sepharose pool four fold with water, and adjusting the pH to 7.5.

four fold with water, and adjusting the pH to 7.5. This mixture was made to 0.7 M ammonium sulfate. Twenty column volume reverse salt gradient was done at 5 mM NaOAC, pH 5.5, 0.2 M to 0M ammonium sulfate.

15 Otherwise, the above steps were identical.

While the present invention has been described in terms of preferred embodiments, it is understood that variations and modifications will occur to those skilled in the art. Therefore, it is intended that the appended claims cover all such equivalent variations which come within the scope of the invention as claimed.

#### CLAIMS

- 1. A method of treating excess weight in a mammal by continuous administration of 1 mg protein/kg body weight/day or less of an OB protein selected from the group consisting of:
  - (a) recombinant methionyl murine OB protein (SEQ. ID. No. 2);
- (b) recombinant methionyl human OB protein (SEQ ID No. 1);
  - (c) the protein of (a) or (b) lacking the methionyl residue at position -1;
- (d) the protein of (a), (b) or (c) lacking a glutamine at position 28; and
  - (e) a chemically modified derivative of (a),(b),(c) or (d).
- 2. A method of claim 1 wherein the 20 chemically modified derivative is a pegylated derivative.
  - 3. A method of claim 2 wherein the pegylated derivative is N-terminally pegylated.
  - 4. A method of claim 1 wherein said continuous administration is accomplished by osmotic pump.
- 5. A DNA sequence according to SEQ ID No. 1.
  - 6. A vector containing a DNA sequence according to claim 5.
- 7. A vector of claim 6 wherein said vector is pCFM1656.

- 8. A DNA sequence according to SEQ ID No. 3.
- 9. A vector containing a DNA sequence 5 according to claim 8.
  - 10. A vector according to claim 9 wherein said vector is pCFM1656.
- 11. A method of refolding partially purified OB protein in a solution obtained from inclusion bodies, said partially purified OB protein selected from the group consisting of:
- (a) recombinant methionyl murine OB protein 15 (SEQ. ID. No. 2);
- (b) recombinant methionyl human OB protein

(SEQ ID No. 1);

- (c) the protein of (a) or (b) lacking the methionyl residue at position -1;
- wherein said refolding is accomplished using N-lauroyl sarcosine.
- 12. A method of claim 11 wherein said sarcosine is used at a concentration of 0.5% 2.0% weight per volume of solution.

#### ABSTRACT

The present invention provides methods and compositions for treating excess weight by administering OB protein in a form for constant supply, at a dosage of less than or equal to about 1 mg protein/kg body weight/day. Compositions and methods used for production of recombinant murine and human OB protein are also provided. Compositions and methods for preparing recombinant murine methionyl OB protein and recombinant human methionyl OB protein, including DNA sequences, vectors, host cells, methods of fermentation, and methods of purification are provided herein.

15

#### SEQUENCE LISTING

(i)	APPLICANT:	Pelleymounter, Ma	ary	Ann
		Hecht, Randy I		
		Mann. Michael B		

- (ii) TITLE OF INVENTION: OB PROTEIN COMPOSITIONS AND METHODS
- (iii) NUMBER OF SEQUENCES: 6
- (iv) CORRESPONDENCE ADDRESS:
  - (A) ADDRESSEE: Amgen Inc.
  - (B) STREET: 1840 Dehavilland Drive
  - (C) CITY: Thousand Oaks
  - (D) STATE: California
  - (E) COUNTRY: U.S.A.
  - (F) ZIP: 91230-1789
- (v) COMPUTER READABLE FORM:
  - (A) MEDIUM TYPE: Floppy disk
  - (B) COMPUTER: IBM PC compatible
  - (C) OPERATING SYSTEM: PC-DOS/MS-DOS
  - (D) SOFTWARE: PatentIn Release #1.0, Version #1.30
- (vi) CURRENT APPLICATION DATA:
  - (A) APPLICATION NUMBER: US 08/474,833
  - (B) FILING DATE: 07-JUN-1995
  - (C) CLASSIFICATION:
- (viii) ATTORNEY/AGENT INFORMATION:
  - (A) NAME: Pessin, Karol M.
  - (C) REFERENCE/DOCKET NUMBER: A-345
- (2) INFORMATION FOR SEQ ID NO:1:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 491 base pairs
    - (B) TYPE: nucleic acid
    - (C) STRANDEDNESS: double
    - (D) TOPOLOGY: linear
  - (ii) MOLECULE TYPE: cDNA
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:

TCTAGATTTG	AGTTTTAACT	TTTAGAAGGA	GGAATAACAT	ATGGTACCGA	TCCAGAAAGT	60
TCAGGACGAC	ACCAAAACCT	TAATTAAAAC	GATCGTTACG	CGTATCAACG	ACATCAGTCA	120
CACCCAGTCG	GTCTCCGCTA	AACAGCGTGT	TACCGGTCTG	GACTTCATCC	CGGGTCTGCA	180
CCCGATCCTA	AGCTTGTCCA	AAATGGACCA	GACCCTGGCT	GTATACCAGC	AGGTGTTAAC	240
CTCCCTGCCG	TCCCAGAACG	TTCTTCAGAT	CGCTAACGAC	CTCGAGAACC	TTCGCGACCT	300
GCTGCACCTG	CTGGCATTCT	CCAAATCCTG	CTCCCTGCCG	CAGACCTCAG	GTCTTCAGAA	360

ACCGGAATCC CTGGACGGGG TCCTGGAAGC ATCCCTGTAC AGCACCGAAG TTGTTGCTCT	420
GTCCCGTCTG CAGGGTTCCC TTCAGGACAT CCTTCAGCAG CTGGACGTTT CTCCGGAATG	480
TTAATGGATC C	491
(2) INFORMATION FOR SEQ ID NO:2:	
<ul> <li>(i) SEQUENCE CHARACTERISTICS:</li> <li>(A) LENGTH: 491 base pairs</li> <li>(B) TYPE: nucleic acid</li> <li>(C) STRANDEDNESS: double</li> <li>(D) TOPOLOGY: linear</li> </ul>	
(ii) MOLECULE TYPE: cDNA	
(xi) SEOUENCE DESCRIPTION: SEO ID NO:2:	

AGATCTAAAC	TCAAAATTGA	AAATCTTCCT	CCTTATTGTA	TACCATGGCT	AGGTCTTTCA	60
AGTCCTGCTG	TGGTTTTGGA	ATTAATTTTG	CTAGCAATGC	GCATAGTTGC	TGTAGTCAGT	120
GTGGGTCAGC	CAGAGGCGAT	TTGTCGCACA	ATGGCCAGAC	CTGAAGTAGG	GCCCAGACGT	180
GGGCTAGGAT	TCGAACAGGT	TTTACCTGGT	CTGGGACCGA	CATATGGTCG	TCCACAATTG	240
GAGGGACGGC	AGGGTCTTGC	AAGAAGTCTA	GCGATTGCTG	GAGCTCTTGG	AAGCGCTGGA	300
CGACGTGGAC	GACCGTAAGA	GGTTTAGGAC	GAGGGACGGC	GTCTGGAGTC	CAGAAGTCTT	360
TGGCCTTAGG	GACCTGCCCC	AGGACCTTCG	TAGGGACATG	TCGTGGCTTC	AACAACGAGA	420
CAGGGCAGAC	GTCCCAAGGG	AAGTCCTGTA	GGAAGTCGTC	GACCTGCAAA	GAGGCCTTAC	480
AATTACCTAG	G					491

### (2) INFORMATION FOR SEQ ID NO:3:

- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 147 amino acids
  - (B) TYPE: amino acid
  - (C) STRANDEDNESS: single
  - (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: protein

#### (xi) SEQUENCE DESCRIPTION: SEQ ID NO:3:

Met Val Pro Ile Gln Lys Val Gln Asp Asp Thr Lys Thr Leu Ile Lys 10 15

Thr Ile Val Thr Arg Ile Asn Asp Ile Ser His Thr Gln Ser Val Ser 25 20 30

Ala Lys Gln Arg Val Thr Gly Leu Asp Phe Ile Pro Gly Leu His Pro 35

Ile Leu Ser Leu Ser Lys Met Asp Gln Thr Leu Ala Val Tyr Gln Gln

		50					55					60				
	Val 65	Leu	Thr	Ser	Leu	Pro 70	Ser	Gln	Asn	Val	Leu 75	Gln	Ile	Ala	Asn	Asp 80
	Leu	Glu	Asn	Leu	Arg 85	Asp	Leu	Leu	His	Leu 90	Leu	Ala	Phe	Ser	Lys 95	Ser
	Cys	Ser	Leu	Pro 100	Gln	Thr	Ser	Gly	Leu 105	Gln	Lys	Pro	Glu	Ser 110	Leu	Asp
	Gly	Val	Leu 115	Glu	Ala	Ser	Leu	Tyr 120	Ser	Thr	G1u	Va1	Val 125	Ala	Leu	Ser
	Arg	Leu 130	Gln	Gly	Ser	Leu	Gln 135	Asp	Ile	Leu	Gln	Gln 140	Leu	Asp	Val	Ser
	Pro 145	Glu	Cys													
(2)	(2) INFORMATION FOR SEQ ID NO:4:															
	<ul> <li>(i) SEQUENCE CHARACTERISTICS:</li> <li>(A) LENGTH: 454 base pairs</li> <li>(B) TYPE: nucleic acid</li> <li>(C) STRANDEDNESS: double</li> <li>(D) TOPOLOGY: linear</li> </ul>															
	(ii)	MOLE	ECULE	E TYF	E: c	DNA										

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:4:

CATATGGTAC	CGATCCAGAA	AGTTCAGGAC	GACACCAAAA	CCTTAATTAA	AACGATCGTT	60
ACGCGTATCA	ACGACATCAG	TCACACCCAG	TCGGTGAGCT	CTAAACAGCG	TGTTACAGGC	120
CTGGACTTCA	TCCCGGGTCT	GCACCCGATC	CTGACCTTGT	CCAAAATGGA	CCAGACCCTG	180
GCTGTATACC	AGCAGATCTT	AACCTCCATG	CCGTCCCGTA	ACGTTCTTCA	GATCTCTAAC	240
GACCTCGAGA	ACCTTCGCGA	CCTGCTGCAC	GTGCTGGCAT	TCTCCAAATC	CTGCCACCTG	300
CCATGGGCTT	CAGGTCTTGA	GACTCTGGAC	TCTCTGGGCG	GGGTCCTGGA	AGCATCCGGT	360
TACAGCACCG	AAGTTGTTGC	TCTGTCCCGT	CTGCAGGGTT	CCCTTCAGGA	CATGCTTTGG	420
CAGCTGGACC	TGTCTCCGGG	TTGTTAATGG	ATCC			454

#### (2) INFORMATION FOR SEQ ID NO:5:

- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 454 base pairs (B) TYPE: nucleic acid

  - (C) STRANDEDNESS: double
    (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: cDNA

	SEQ	UENCI	E DES	CRI	PTIO	N: S	EQ I	D NO	:5:							
GTATACCA	TG G	CTAG	GTCTT	' TC	AAGT	CCTG	CTG	rggt:	$\Gamma T T$	GGAA'	raati	T T	rgct.	AGCA	A	60
TGCGCATA	GT T	GCTG:	PAGTO	: AG	rgtg	GGTC	AGC	CACTO	CGA	GATT:	rgtco	C A	CAAT(	GTCC	G	120
GACCTGAA	GT A	GGGC	CCAGA	. CG	rggg(	CTAG	GAC'	rggaz	ACA	GGTT.	rtacc	T G	GTCT(	GGGA	C	180
CGACATAT	GG T	CGTC	ΓAGAA	TTC	GGAG(	GTAC	GGC	AGGG(	CAT	TGCA	AGAAG	T C'	raga(	GATT(	G	240
CTGGAGCT	CT T	GGAA(	GCGCT	' GGZ	ACGA	CGTG	CAC	GACC	TA.	AGAG(	STTTA	G G	ACGG'	rgga(	C	300
GGTACCCG	AA G	TCCAC	GAACT	CTC	GAGAC	CCTG	AGA	GACCO	CGC	CCCA	GGACC	T T	CGTA	GGCCZ	A	360
ATGTCGTG	GC T	TCAA(	CAACG	AGA	ACAGO	GCA	GAC	GTCC	CAA	GGGAZ	AGTCC	T G	racg <i>i</i>	AAAC(	C	420
GTCGACCT	GG A	CAGA(	GCCC	AAC	CAATT	TACC	TAGO	ב כ								454
(2) INFO	RMAT	ION E	FOR S	EQ ]	ED NO	):6:										
(i)	(A	TYF) STF	IGTH:	147 mino DNES	7 ami 5 aci 55: s	ino a ld singl	acids	S								
(ii)	MOL:	ECULE	E TYP	E: p	prote	ein										
(xi)	SEQ	UENCE	E DES	CRIE	OITC	N: SE	EQ II	O NO:	6:							
Met 1	Val	Dro	<del>-</del> 1 -	Cl ~	_	_										
		FLO	TTE	5	Lys	Val	Gln	Asp	Asp 10	Thr	Lys	Thr	Leu	Ile 15	Lys	
Thr	Ile	Val		5					10					15		
Thr Ser Ly:		Val	Thr 20	5 Arg	Ile	Asn	Asp	Ile 25	10 Ser	His	Thr Gly	Gln	Ser 30	15 Val	Ser	
Ser Ly:	s Gl:	Val n Arg	Thr 20 y Val	5 Arg Thi	Ile Gly	Asn / Let	Asp 1 Asp 40	Ile 25 Phe	10 Ser	His e Pro	Thr Gly	Gln Let 45	Ser 30 1 His	15 Val s Pro	Ser	
Ser Ly: Ile	s Gl: Leu 50	Val n Arg 35	Thr 20 Val Leu	5 Arg Thi	Ile Gly Lys	Asn Let Met 55	Asp 1 Asp 40 Asp	Ile 25 Phe Gln	10 Ser	His e Pro	Thr Gly Ala 60	Gln Lei 45 Val	Ser 30 1 His	15 Val S Pro	Ser O Gln	
Ser Lys Ile Ile 65	Leu 50	Val n Arg 35 Thr	Thr 20 Val Leu Ser	5 Arg Thi Ser Met	Ile Gly Lys Pro 70	Asn Lew Met 55 Ser	Asp 40 Asp Arg	Ile 25 Phe Gln Asn	10 Ser Il Thr	His e Pro Leu Leu 75	Thr Gly Ala 60 Gln	Gln Lev 45 Val	Ser 30 His Tyr	15 Val S Pro Gln Asn	Ser Gln Asp 80	
Ser Lys Ile Ile 65 Leu	Leu 50 Leu Glu	Val n Arg 35 Thr	Thr 20 Val Leu Ser	5 Arg Thr Ser Met Arg 85	Ile Gly Lys Pro 70 Asp	Asn Let S5 Ser Leu	Asp  Asp  Asp  Arg	Ile 25 Phe Gln Asn	10 Ser Il Thr Val Val 90	His e Pro Leu 75 Leu	Thr Gly Ala 60 Gln Ala	Gln Lev 45 Val Ile Phe	Ser 30 His Tyr Ser	15 Val S Pro Gln Asn Lys 95	Ser Gln Asp 80 Ser	
Ser Lys Ile Ile 65 Leu Cys	Leu 50 Leu Glu	Val n Arg 35 Thr Thr	Thr 20 Val Leu Ser Leu Pro 100	5 Arg Thr Ser Met Arg 85 Trp	Ile Gly Lys Pro 70 Asp	Asn Let Ser	Asp 1 Asp 40 Asp Arg Leu	Ile 25 Phe Gln Asn His Leu 105	10 Ser Il Thr Val Val 90 Glu	His e Pro Leu 75 Leu	Thr OGLY Ala 60 Gln Ala Leu Val	Gln Lev 45 Val Ile Phe	Ser 30 His Tyr Ser Ser 110	Val Val Gln Asn Lys 95 Leu	Ser  Gln  Asp 80  Ser  Gly	
Ser Lys Ile Ile 65 Leu Cys Gly	Leu 50 Leu Glu His	Val n Arg 35 Thr Asn Leu Leu	Thr 20 Val Leu Ser Leu Pro 100 Glu	5 Arg Thr Ser Met Arg 85 Trp	Ile CGly Lys Pro 70 Asp Ala Ser	Asn Let Ser Gly	Asp  1 Asp 40 Asp  Arg  Leu Gly Tyr 120	Ile 25 Phe Gln Asn His Leu 105 Ser	10 Ser Il Thr Val 90 Glu Thr	His e Pro Leu 75 Leu Thr	Thr OGLY Ala 60 Gln Ala Leu Val	Gln Lev 45 Val Ile Phe Asp Val 125	Ser 30 His Tyr Ser Ser 110 Ala	Val Val Gln Asn Lys 95 Leu Leu	Ser  Gln  Asp 80  Ser  Gly  Ser	

# DECLARATION AND POWER OF ATTORNEY

As a below named inventor, I hereby declare that:

My residence, post office address and citizenship are as stated below next to my name.

I believe I am the original, first, and sole inventor (if only one name is listed below) or a joint inventor (if plural names are listed below) of the invention entitled

# OB PROTEIN COMPOSITIONS AND METHODS

which is described and claimed in the specification which	) <b>:</b>
X is attached hereto.	
was filed on	
as Application Serial No.:	
and was amended on (if app	licable)
I hereby state that I have reviewed and understand the including the claims, as amended by any amendment ref	he contents of the above identified specification, ferred to above.
I acknowledge the duty to disclose information which is accordance with Title 37, Code of Federal Regulations,	s material to the examination of this application in §1.56(a).
Power of Attorney: As a named inventor, I hereby approsecute this application and transact all business therewith:	point the following attorney(s) and/or agent(s) to in the Patent and Trademark Office connected
Ron K. Levy, Registration No.: 31,539, Steven M. Odre Registration No. 34,899, said attorney(s)/agent(s) to he the power to revoke any power herein granted.	Registration No.: 29,094, and Karol M. Pessin, ave in addition full power of revocation, including
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I hereby declare that all statements made herein of my own knowledge are true and that all statements made on information and belief are believed to be true; and further that these statements were made with the knowledge that willful false statements and the like so made are punishable by fine or imprisonment, or both, under section 1001 of Title 18 of the United States Code, and that such willful false statements may jeopardize the validity of the application or any patent issuing thereon.

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Applicants: Pelleymounter et al.

Serial No.: 08/920,608

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For:

**OB Protein Compositions and Methods** 

Docket No.: A-345A

# ASSOCIATE POWER OF ATTORNEY

Assistant Commissioner for Patents Washington, D.C. 20231

Sir:

Please recognize <u>Joan D. Eggert</u>, Registration No.: <u>32.980</u>, as associate attorney in this application, with full power to prosecute the application, to make alterations and amendments therein, and to transact such other business in the Office in connection therewith as may be necessary.

Respectfully submitted,

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